

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/586,403  
Source: IFWP  
Date Processed by STIC: 07/26/2006

# ***ENTERED***



IFWP

## RAW SEQUENCE LISTING

DATE: 07/26/2006

PATENT APPLICATION: US/10/586,403

TIME: 14:18:54

Input Set : A:\00786.455003.SEQLIST.TXT

Output Set: N:\CRF4\07262006\J586403.raw

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4 <110> APPLICANT: RAHME, Laurence
5     DEZIEL, Eric
6     LEPINE, Francois
7     TOMPKINS, Ronald G.
8     XIAO, Gaoping
10 <120> TITLE OF INVENTION: Methods For Identifying Candidate
11     Compounds For Treating, Reducing, or Preventing Pathogenic
12     Infections
14 <130> FILE REFERENCE: 00786/455003
C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/586,403
C--> 16 <141> CURRENT FILING DATE: 2006-07-18
16 <150> PRIOR APPLICATION NUMBER: PCT/US05/02174
17 <151> PRIOR FILING DATE: 2005-01-21
19 <150> PRIOR APPLICATION NUMBER: US 60/538,361
20 <151> PRIOR FILING DATE: 2004-01-22
22 <150> PRIOR APPLICATION NUMBER: US 60/538,278
23 <151> PRIOR FILING DATE: 2004-01-21
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31 <212> TYPE: DNA
32 <213> ORGANISM: Pseudomonas aeruginosa
34 <400> SEQUENCE: 1
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37 tcccaggcca gccaactggc ccgcctgctc aagcccggcg atcgcggtgt gctggcggtt 180
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43 ctgcaataca cctcgggttc caccggggcg cccaaggggg tgatgcacag cctgcgcaac 540
44 acgctcgggt tctgcggggc gttcgctacg gagtgtgctg cattgcaggc gggagaccgg 600
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48 ctgcgtccgc aggccaggga gctggtgagc agcgtgcgcc tggcgttttc cgccggctcg 840
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50 atcggggcta ccgaggtcgg ccatgtgttc ctgcccaacc gcccgggcca ggcgcggtgc 960
51 gacagcaccg ggctgcggtt gcctggctat gagtgcgggc tgggtggaccg cgaaggacac 1020
52 actatcgagg aagcggggcg gcaaggcgtg ctggttggtg gtggcccagg gctgagtcgg 1080
53 ggttactggc gggccagcga agagcagcag gcgcgcttcg caggtggctg gtaccgcacc 1140

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55 ctgttcaagg tgaatggccg ctgggtgggt cgcaccagg tcgagcagg gatctgccgt 1260
56 catctgccgg aagtgagcga ggcggttctg gttcctacct gccggtgca cgacggcttg 1320
57 cgtccgaccc tgttcgtcac cctggccact ccgctggacg acaaccagat cctgctggcg 1380
58 cagcgcatcg accagcatct cgccgaacag attccctcgc acatgctgcc cagccaattg 1440
59 catgtgctgc cggccttgcc gcgcaacgac aacggcaagt tggcgcgcg cgagctgcgc 1500
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64 <212> TYPE: DNA
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70 gcggttcgcg aagccctgga cgaggcgcg gtcaagcccg aggagatcga cctgatcgtc 180
71 ggcctcgccc tgtctccgga ccatctgacg gagaaccgcg acatcatggc gccgaagatc 240
72 ggccatccgt tgcagaaggc cctcggcgcg aatcgcgcg atgtcttcga cctcaccgac 300
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85 <211> LENGTH: 1047
86 <212> TYPE: DNA
87 <213> ORGANISM: Pseudomonas aeruginosa
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92 gtcaatcggc ggggtatatt cgaccgcgg aacggcgaga acgagttcag cctggtggtc 180
93 cgggcgcgag agcgctgct gcgtagcagc gataccgcgc cggatagcgt ggacatgctg 240
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95 ggacgtctct acccgcgcat ggccaacgtg ctgtccaagc agctcgccct gagtccggcg 360
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99 ctgctgaccc gcggcgacga tgacagctgc gacctgctgg cttcggccga acacagcgac 600
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101 gccaagcgcg ggttttattt ctggtgttc agcgacggcc agaacaagat ggccagcttc 720
102 gttccgacca acgtgccgat cgcgatgcgc cggcgcttg aaaaggcccg cctgggcagc 780
103 gatgacatcg attattctgt cttccaccag ccagcgccgt tctggtcaa ggccggggcc 840
104 gagggcatcg gtgcccgtcc tgagcagtac caactgacga tgggcgatac cggcgtgatg 900
105 atctcgtttt ccatccgta caccctgatg accggcctgc gcgagggcaa gatccgcccc 960
106 ggcgatcgta tcgtcatggc cggcgacgcc actggctggg ggttcgcccgc ccaggctctg 1020

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114 <400> SEQUENCE: 4
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116 catgacctgg tagggcgcat caatacgtcg gacgagttca tcgtcgaacg taccggcggtg 120
117 cgcacccgct atcacgtcga gccggaacag gcggtcagcg cgctgatggt gccggcgggcg 180
118 cgccaggcca tcgaggctgc cgggctgctg ccggaggaca tcgacctgtt gctggtgaac 240
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120 ctgcggcaca tcccgggtact ggatatccgg gcacagtga gcgggttgc gtacggcttg 360
121 cagatggctc gcgggcagat cctcgccggg ctggcacggc atgtcctggt ggtctgcggc 420
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128 aacctgcgca tctcgatgc ggtgcaggag caactgggca tccccagca caagttcgcg 840
129 gtgacctgg atcgtctggg caacatggct tcggcctcga ccccggtcac gctggcgatg 900
130 ttctggccgg acatccagcc gggacagcgg gtgctggtcc tgacctacgg ctccggcgcg 960
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141 atcagccggg atgcgcaatt ggtttggcg gacctgtgcc gctgggtcgc cgacctgcc 180
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144 aagtcggaaa gcgcgggtgc ggtgggtcgag cgcttgaacc ggcaactgtt gcgtgcggag 360
145 cagcggttgc ccgaggcctg tgctggggac gctctgccgg ttcgcgcggg ggccgacggc 420
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152 ctgagcgagg agctgcaccg cgctgggggt gggcgagagc tcgacttcct gcccgcgcaa 840
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154 gactga 906
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157 <211> LENGTH: 1149
158 <212> TYPE: DNA
159 <213> ORGANISM: Pseudomonas aeruginosa
161 <400> SEQUENCE: 6

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164 ggcatacccc tggcgagcaa tgcgttgacg gcgttggtcca gcacctgga tctcgaccgg 180
165 ctgttccgcc gtggcatgcc gttggccggc atcaacgtat acgcccacga cggttcgatg 240
166 ctgatgtcga tgccttcag tctgggtggg aattcccggc gggcctggc gttgcagcgc 300
167 cacgaactgc atgcggcgct actggagggg ctggatgagt cgcgcattcg ggtcgggggtc 360
168 tccatcgtgc agatcctcga cggactcgac cacgaacgcy tgacctgag cgacggcact 420
169 gtccacgact gttcgtggt ggtcgggtgc gatggcattc gttcgagcgt gcgacgttat 480
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171 catcggtctg aggacgccga gctggcgggg gaggtctggg ggcacggcaa gcgcctcggc 600
172 ttcataccaga tcagcccggc cgagatgtat gtctacgcga ccctgaaggc gcgcccggag 660
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175 aacgacctcg aggagtggc cggcgccctc tggtgccgcy gacgggtagt gctgatcggt 840
176 gacgcccac acgccatgac gccgaacctg gggcaggggc cggccatggc cctggaggac 900
177 gccttccctgc tggcgcgctt gtggtgctgc gcgcgcgcy ccgagacgct gatcctgttc 960
178 cagcagcaac gcgaggcgcy gatcgagttc atcaggaagc aatcctggat cgtcggccgc 1020
179 cttgggtcagt gggaatcgcc ctggagcgtc tggctgagga atacctcgt tcgcctggtg 1080
180 ccgaatgcca gtgcgaggcg cctccaccag cgtcttttca ccggtgtcgg tgagatggcc 1140
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184 &lt;211&gt; LENGTH: 1197

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186 &lt;213&gt; ORGANISM: Pseudomonas aeruginosa

188 &lt;400&gt; SEQUENCE: 7

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191 cgcgcgatca acggcgccga cctgctcaag ccggccggca tccgggtggt cgaggcgcc 180
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195 accgtcgaga tgtgttcga gaccgcata gaagcgggtc agcgcgacga gcgccacgcy 420
196 atcgaccagg tgcgcctgaa cgacggccgc gtgctgcgcy cgcgggtggt ggtgggagcc 480
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198 taccctgcgc cgatgctggt cggcaccttc gccctggcgc cctgcgtggc cgagcgcaac 600
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212 &lt;212&gt; TYPE: PRT

213 &lt;213&gt; ORGANISM: Pseudomonas aeruginosa

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220 Leu Gln Cys Arg Thr Tyr Ile Leu Ser Gln Ala Ser Gln Leu Ala Arg
221 35 40 45
222 Leu Leu Lys Pro Gly Asp Arg Val Val Leu Ala Leu Asn Asp Ser Pro
223 50 55 60
224 Ser Leu Ala Cys Leu Phe Leu Ala Cys Ile Ala Val Gly Ala Ile Pro
225 65 70 75 80
226 Ala Val Ile Asn Pro Lys Ser Arg Glu Gln Ala Leu Ala Asp Ile Ala
227 85 90 95
228 Ala Asp Cys Gln Ala Ser Leu Val Val Arg Glu Ala Asp Ala Pro Ser
229 100 105 110
230 Leu Ser Gly Pro Leu Ala Pro Leu Thr Leu Arg Ala Ala Ala Gly Arg
231 115 120 125
232 Pro Leu Leu Asp Asp Phe Ser Leu Asp Ala Leu Val Gly Pro Ala Asp
233 130 135 140
234 Leu Asp Trp Ser Ala Phe His Arg Gln Asp Pro Ala Ala Ala Cys Phe
235 145 150 155 160
236 Leu Gln Tyr Thr Ser Gly Ser Thr Gly Ala Pro Lys Gly Val Met His
237 165 170 175
238 Ser Leu Arg Asn Thr Leu Gly Phe Cys Arg Ala Phe Ala Thr Glu Leu
239 180 185 190
240 Leu Ala Leu Gln Ala Gly Asp Arg Leu Tyr Ser Ile Pro Lys Met Phe
241 195 200 205
242 Phe Gly Tyr Gly Met Gly Asn Ser Leu Phe Phe Pro Trp Phe Ser Gly
243 210 215 220
244 Ala Ser Ala Leu Leu Asp Thr Trp Pro Ser Pro Glu Arg Val Leu
245 225 230 235 240
246 Glu Asn Leu Val Ala Phe Arg Pro Arg Val Leu Phe Gly Val Pro Ala
247 245 250 255
248 Ile Tyr Ala Ser Leu Arg Pro Gln Ala Arg Glu Leu Leu Ser Ser Val
249 260 265 270
250 Arg Leu Ala Phe Ser Ala Gly Ser Pro Leu Pro Arg Gly Glu Phe Glu
251 275 280 285
252 Phe Trp Ala Ala His Gly Leu Glu Ile Cys Asp Gly Ile Gly Ala Thr
253 290 295 300
254 Glu Val Gly His Val Phe Leu Ala Asn Arg Pro Gly Gln Ala Arg Ala
255 305 310 315 320
256 Asp Ser Thr Gly Leu Pro Leu Pro Gly Tyr Glu Cys Arg Leu Val Asp
257 325 330 335
258 Arg Glu Gly His Thr Ile Glu Glu Ala Gly Arg Gln Gly Val Leu Leu
259 340 345 350
260 Val Arg Gly Pro Gly Leu Ser Pro Gly Tyr Trp Arg Ala Ser Glu Glu
261 355 360 365
262 Gln Gln Ala Arg Phe Ala Gly Gly Trp Tyr Arg Thr Gly Asp Leu Phe
263 370 375 380

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Input Set : A:\00786.455003.SEQLIST.TXT

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L:16 M:270 C: Current Application Number differs, Replaced Current Application No

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date